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Asp Pro Glu Asn Val His Tyr Ile Met Ser Ala Asn Phe Gln Asn Phe 85 90 95

Pro Lys Gly Pro Lys Phe Arg Glu Ile Phe Asp Val Leu Gly Asp Gly 100 105 110

Ile Phe Asn Ala Asp Ser Glu Ser Trp Arg Asp Gln Arg Arg Val Ala

115 120 125

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420 425 430

Ile Ser Asp Arg Gly Ser Ile Lys His Glu Pro Ser Tyr Lys Phe Leu 440 Ala Phe Asn Ala Gly Pro Arg Thr Cys Leu Gly Lys Asp Val Ala Phe Ala Gln Val Lys Ala Val Ala Ala Thr Leu Ile His Asn Tyr Gln Val 470 475 His Val Ala Asp Gly His Arg Val Leu Pro Asn Cys Ser Ile Ile Leu 485 Tyr Met Arg Asn Gly Leu Lys Val Arg Val Ala Asn Arg Trp Ser Ala 500 505 Lys Lys Asn 515 <210> 57 <211> 1494 <212> DNA <213> Sesamum indicum <220> <223> SiP168 <400> 57 atggatctac tactttccct tgttctccta ctctgttctg cagcatgcat ttggtttctc 60 cgggtggtcc tgaaacccaa tccagggccc cggaaatcag ccaatctccc tccagggcca 120 aaacctcttc ccataatcgg caacattctt gagcttggtg agaaacccca ccaatctctc 180 gccaaactct ccaaaaccta cgggcccctg atgcgtctca agctgggaac catgacaaca 240 gttgttgtat cctccccgga aatctccagg atcgtgctgc aacaatatga ccaagttttc 300 tccagccgaa cacacgcaga tgccatccga gcacttgacc accacaaaca ttccgtcgcc 360 tggataccgg cggacaatca gtggcggaaa atccgtaaac tgtgcaaaqa qaaqatqttt 420 tcgggccaaa agcttgatgc gaaccagggc ctgaggaggg agaagttgcg taatttgcaa 480 gactatgtga atgaatgctg cgttagtggc caggtcgtgg atattggtgt agctgccttt 540 acgacgaccc ttaatctgat atcggccact cttttctcgg tggattttgc tgattttggt 600 tctggttcgt ctcaagagct taaggatgtt atgagcggga tagcgtctat catcggccga 660 ccaaattttg ctgattgttt ccctcttctt cggctggttg atccacaggg catcttccgc 720 cagaccacgt tacatttcaa caagtgtttt aagatctttg atgaaattat ccgtcaaagg 780 ctacagacca atgattcggg gacgaaaagt gacatgctga aagagcttct tgaaatcaac 840 cagaaagatg agtctgaatt gagctttgac gagatcaagc atttactcct ggatctactt 900 gtcgcaggaa cggacacaac ttcagttaca gtggaatggg caatgacgga gctagtgcgc 960 caccctgaga aaatgtcgaa agccagaaat gagttaagaa atgtggtggg actgaataaa 1020 gaaattcaag aatcagacat ctcaagactc ccttacctac gagcagtggt gaaagaaagt 1080 ttcaggcttc accctgcaac tcctttatcg gtacctcaca aggccgacga ggaagcagaa 1140 atcaatggct atatagtccc taaaggagca caagttctca tgaacgtgtg ggccatcggc 1200 agagattcaa gcatatggag gaaccctgat gtattcatgc ccgagaggtt cttggagaca 1260 gaaattgatg teegtggeea acaettegag etgetteett ttggeggggg gaggaggatt 1320 tgcgtggggc tgccgttagc ctatcgtatg atccatctcg tgcttgccac tttcataagc 1380 gactatgatt ggaaacttga aggagggctg aaaactgaag aaatggacat gagtgaaaag 1440

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<213> Sesamum indicum

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Ile Leu Glu Leu Gly Glu Lys Pro His Gln Ser Leu Ala Lys Leu Ser 50 55 60

Lys Thr Tyr Gly Pro Leu Met Arg Leu Lys Leu Gly Thr Met Thr Thr 65 70 75 80

Val Val Val Ser Ser Pro Glu Ile Ser Arg Ile Val Leu Gln Gln Tyr 85 90 95

Asp Gln Val Phe Ser Ser Arg Thr His Ala Asp Ala Ile Arg Ala Leu 100 105 110

Asp His His Lys His Ser Val Ala Trp Ile Pro Ala Asp Asn Gln Trp 115 120 125

Arg Lys Ile Arg Lys Leu Cys Lys Glu Lys Met Phe Ser Gly Gln Lys 130 135 140

Leu Asp Ala Asn Gln Gly Leu Arg Arg Glu Lys Leu Arg Asn Leu Gln 145 150 155 160

Asp Tyr Val Asn Glu Cys Cys Val Ser Gly Gln Val Val Asp Ile Gly
165 170 175

Val Ala Ala Phe Thr Thr Leu Asn Leu Ile Ser Ala Thr Leu Phe 180 185 190

Ser Val Asp Phe Ala Asp Phe Gly Ser Gly Ser Ser Gln Glu Leu Lys 195 200 205

Asp Val Met Ser Gly Ile Ala Ser Ile Ile Gly Arg Pro Asn Phe Ala 210 215 220

Asp Cys Phe Pro Leu Leu Arg Leu Val Asp Pro Gln Gly Ile Phe Arg 225 230 235 240

Gln Thr Thr Leu His Phe Asn Lys Cys Phe Lys Ile Phe Asp Glu Ile
245 250 255

Ile Arg Gln Arg Leu Gln Thr Asn Asp Ser Gly Thr Lys Ser Asp Met 260 265 270

Leu Lys Glu Leu Leu Glu Ile Asn Gln Lys Asp Glu Ser Glu Leu Ser 275 280 285

Phe Asp Glu Ile Lys His Leu Leu Leu Asp Leu Leu Val Ala Gly Thr 290 295 300

Asp Thr Thr Ser Val Thr Val Glu Trp Ala Met Thr Glu Leu Val Arg 305 310 315 320

His Pro Glu Lys Met Ser Lys Ala Arg Asn Glu Leu Arg Asn Val Val 325 330 335

Gly Leu Asn Lys Glu Ile Gln Glu Ser Asp Ile Ser Arg Leu Pro Tyr 340 345 350

Leu Arg Ala Val Val Lys Glu Ser Phe Arg Leu His Pro Ala Thr Pro 355 360 365

Leu Ser Val Pro His Lys Ala Asp Glu Glu Ala Glu Ile Asn Gly Tyr 370 375 380

Ile Val Pro Lys Gly Ala Gln Val Leu Met Asn Val Trp Ala Ile Gly 385 390 395 400

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Phe Leu Glu Thr Glu Ile Asp Val Arg Gly Gln His Phe Glu Leu Leu 420 425 430

Pro Phe Gly Gly Arg Arg Ile Cys Val Gly Leu Pro Leu Ala Tyr 435 440 445

Arg Met Ile His Leu Val Leu Ala Thr Phe Ile Ser Asp Tyr Asp Trp 450 455 460

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Val Phe Asn Gln Leu Leu Asn Phe Tyr Arg Leu His Asp Tyr Met Ala
Asp Leu Ala Gly Lys Tyr Lys Thr Tyr Arg Leu Ile Ala Pro Phe Arg
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Thr Asn Phe Glu Ser Tyr Gly Lys Gly Pro Tyr Asn Cys Ser Ile Leu
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Gly Asp Leu Phe Gly Glu Gly Ile Phe Ala Ile Asp Gly His Lys Trp 115 120 Arg Glu Gln Arg Lys Val Ser Ser Leu Glu Phe Ser Thr Arg Val Leu Arg Asp Tyr Ser Ser Ile Val Phe Arg Lys Asn Ala Val Arg Leu Ala Lys Ile Leu Ser Gly Ala Ala Thr Ser Asn Gln Pro Val Asp Ile Gln Asp Leu Phe Met Lys Ser Thr Phe Asp Ser Ile Ser Glu Val Ala Leu Gly Val Glu Leu Asp Ser Leu Gly Gly Ser Asn Glu Glu Gly Ala Lys 200 Phe Ser Ile Ala Ala Asp Asp Val Ser Met Arg Thr Leu Trp Arg Tyr 210 215 Val Asp Val Leu Trp Lys Leu Lys Arg Ala Leu Asn Val Gly Ser Glu 230 Ala Lys Leu Lys Lys Ser Leu Gln Val Val Asp Glu Phe Val Tyr Lys 245 250 Leu Ile His Ser Arg Thr Gln Gln Met Asn Met Pro Gly Asn Asp Ser Val Met Gln Leu Lys Lys Asp Asp Ile Leu Ser Arg Phe Leu Gln Leu 280 Thr Glu Ala Thr Pro Lys Tyr Leu Arg Asp Ile Thr Ile Ser Phe Ile 290 Val Ala Gly Lys Asp Thr Thr Ala Thr Thr Leu Ser Trp Phe Ile Tyr Met Leu Cys Lys Tyr Pro His Val Gln Glu Lys Val Glu Gln Glu Ile 325 Lys Asp Ala Thr Gly Cys Lys Glu Val Ala Asp Ile Ser Glu Phe Ser Ala Cys Val Thr Glu Glu Ala Leu Gly Lys Met His Tyr Leu His Ala 360 Ala Leu Thr Glu Thr Leu Arg Ile Tyr Pro Ala Val Ala Val Asp Ala 370 Lys Gln Cys Leu Cys Asp Asp Ile Met Pro Asp Gly Phe Ser Val Lys Lys Gly Asp Met Val Ala Tyr Gln Pro Tyr Ala Met Gly Arg Met Lys

				405					410					415		
Ser	Ile	Trp	Gly 420	Asn	Asp	Ala	Glu	Glu 425	Phe	Lys	Pro	Glu	Arg 430	Trp	Leu	
Asp	Lys	Asn 435	Gly	Cys	Phe	Gln	Gln 440	Ala	Ser	Pro	Phe	Lys 445	Phe	Thr	Ala	
	Gln 450	Ala	Gly	Pro	Arg	Leu 455	Cys	Leu	Gly	Lys	Glu 460	Phe	Ala	Tyr	Arg	
Gln 465	Met	Lys	Ile	Phe	Ser 470	Ala	Ile	Leu	Leu	Arg 475	Phe	Phe	Thr	Met	Lys 480	
Leu	Ser	Asp	Glu	Arg 485	Lys	Thr	Val	Asn	Tyr 490	Arg	Pro	Met	Leu	Thr 495	Leu	
Leu	Ile	Asp	Gly 500	Gly	Leu	Ile	Val	Arg 505	Pro	Phe	His	Arg	Met 510	Asp	Glu	
Lys	Thr	Ala 515														
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<210	> 62	2														
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<213	> Al	LIII	ıcıa.	l Sec	queno	:e										
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35 40 45

Val His Leu Met Lys Asn Leu Leu His Arg Thr Leu Tyr Asp Phe Ser 50 55 60

Gln Lys Leu Gly Pro Ile Phe Ser Leu Arg Phe Gly Thr Arg Leu Val 65 70 75 80

Val Val Val Ser Ser Ser Ser Leu Val Glu Glu Cys Phe Thr Lys Tyr 85 90 95

Asp Ile Val Leu Ala Asn Arg Pro Gln Pro Ser Val Asp Arg Arg Ser 100 105 110

Leu Gly Phe Ser Thr Thr Ser Val Ile Gly Ala Pro Tyr Gly Asp His 115 120 125

Trp Arg Asn Leu Arg Lys Leu Cys Asp Leu Glu Val Phe Ala Pro Thr 130 135 140

Arg Leu Ala Ser Phe Leu Ser Ile Arg Leu Asp Glu Arg Asp Arg Met 145 150 155 160

Ile Ser Ser Leu Tyr Lys Ile Ser Ser Ala Gly Phe Ala Lys Val Asn 165 170 175

Leu Glu Thr Lys Ile Val Glu Leu Thr Phe Asn Asn Ile Met Arg Met 180 185 190

Val Ala Gly Lys Arg Tyr Tyr Gly Glu Glu Ala Glu Asp Asp Glu Glu
195 200 205

Ala Lys Arg Phe Arg Asp Leu Thr Lys Glu Ala Leu Glu Leu Thr Ser 210 215 220

Ala Ser Asn Pro Gly Glu Ile Phe Pro Ile Leu Arg Trp Leu Gly Phe 225 230 235 240

Asn Gly Leu Glu Lys Lys Leu Ala Val His Ala Arg Lys Thr Asp Glu 245 250 255

Phe Met Gln Gly Leu Leu Asp Glu His Arg Arg Gly Glu Arg Gln Asn 260 265 270

Thr Met Val Asp His Leu Leu Ser Leu Gln Glu Ser Gln Pro Glu Tyr 275 280 285

Tyr Thr Asp Glu Ile Ile Thr Gly Leu Ile Val Ala Leu Ile Ile Ala 290 295 300

Gly Thr Asp Ala Ser Val Val Thr Thr Glu Trp Ala Met Ser Leu Ile 305 310 315 320

Leu Asn His Pro Gln Val Leu Glu Lys Ala Arg Lys Glu Leu Asp Thr 325 330 335

Leu Val Gly His Glu Arg Met Val Asp Glu His Asp Leu Pro Lys Leu 340 345 350

Arg Tyr Leu His Cys Ile Val Leu Glu Thr Leu Arg Leu Phe Pro Ser 355 360 365

Val Pro Thr Leu Val Pro His Glu Pro Ser Glu Asp Cys Lys Ile Gly 370 380

Gly Tyr Asn Val Pro Lys Gly Thr Met Ile Leu Val Asn Ala Trp Ala 385 390 395 400

Ile His Arg Asp Pro Lys Val Trp Asp Asp Pro Leu Ser Phe Lys Pro 405 410 415

Asp Arg Phe Glu Thr Met Glu Val Glu Thr His Lys Leu Leu Pro Phe 420 425 430

Gly Met Gly Arg Arg Ala Cys Pro Gly Ala Gly Leu Ala Gln Lys Phe
435 440 445

Val Gly Leu Ala Leu Gly Ser Leu Ile Gln Cys Phe Glu Trp Glu Arg 450 455 460

Met Ser Ala Glu Lys Ile Asp Leu Asn Glu Gly Ser Gly Ile Thr Leu 465 470 475 480

Pro Lys Ala Lys Thr Leu Glu Ala Met Cys Lys Pro Arg His Ile Met 485 490 495

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Leu Gly Ala Ser Thr Ile Ser Val Ile Gly Ala Pro Tyr Gly Asp His 115 120 125

Trp Arg Asn Leu Arg Lys Leu Cys Asp Leu Glu Val Phe Ala Pro Thr 130 135 140

Arg Leu Ala Ser Phe Leu Ser Ile Arg Arg Asp Glu Arg Asp Arg Met 145 150 155 160

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Leu Gly Phe Ser Thr Thr Ser Val IIe Gly Ala Pro Tyr Gly Asp His 115 120 125

Trp Arg Asn Leu Arg Lys Leu Cys Asp Leu Glu Val Phe Ala Pro Thr $130\,$

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Ile Ser Ala Leu Tyr Lys Ile Ser Ser Ala Gly Phe Ala Lys Val Asn 165 170 175 Francisco

Leu Glu Ala Lys Ile Val Glu Leu Thr Phe Asn Asn Ile Met Arg Met 180

Val Ala Ala Lys Arg Tyr Tyr Gly Glu Glu Ala Glu Asp Asp Glu Glu 195 200 205

Ala Lys Arg Phe Arg Asp Leu Thr Lys Glu Ala Leu Glu Leu Thr Ser 210 220

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Arg Pro Pro Gly Pro Pro Gly Leu Pro Phe Leu Gly Asn Leu Leu Gln

est cheans to type plant personals than Tyr Asn Pro Ser Asp Leu His Leu Arg Leu Thr Lys Leu Ser Glu Lys 50 60

Tyr Gly Pro Leu Met Tyr Met Thr Phe Val Gly Lys Pro Val Val 65 70 75 80 90 Jan 208

to and the contract of the angle of the contract of the contra lle Ser Ser Ala Arg Val Ala Lys Glu Ala Leu Lys Tyr Asn Asp Leu

Ala Phe Ser Ser Arg Pro Ser Thr Ile Ala Ser Arg Lys Val Ala Tyr

Asn Asn Ser Aspalle: Ser Meto Ser Pro TyroThroGluaTyroTrp Arg Gluadano (1881) 15 to 188 (1881) 16 to 188 (18

Leu Arg Lys Met Val Val Leu Arg Leu Phe Thr Val Lys Gln Val Asn $13\overline{0}$ the solution of the 135 galax constants of 140 matrix of the expectation of 1 . The 1

Ser Phe ArgaPro Ala Arg Glu Glu Glu Val Ala Arg Met Val Lys Glus and Arga Met Val Lys Glus Arga Met Va

145 Cold 150 Man Ala His Gln Pro Val Asn Ile Asn Glu Ile

Ala Leu Ser Leu Ser Ser Arg Met Ile Ser Arg Phe Ala Leu Gly Lys 180 - 180 - 190 - 1

Arg Tyr AspaGlu Glu Asn Gly Pro Glu Lys Arg Arg Phe Asp Arg Ile 76 (195) (Maring 1809) (Maring R. 200) (Maring Maring Maring 205), Maring Maring (Maring Maring Mari

Leu Gln Leu Leu Gln Leu Val Ser Val Glu Ile Phe Phe Gly Asp Tyr $210\,$ M , gas at longer trans215, that is long along $r=220\,
m ggs$ (). We show that

March Street, State Street, Callette. Ser Pro Trp Leu Gly Trp Ile Asp Arg Leu Cys Gly Lys Val Ser Gln 225 235 240

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Ala Glu His Leu Ser Pro Asn Arg Pro Glu Ser Met Asn Gly Asp Ile 260 265 270

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His Val Ala Asp Gly His Arg Vale Leu Pro Asn Cys Ser Ile Ile Leu 485 mg 490 mg 495 mg

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lle Arg Gln Arg Leu Gln Thr Asn Asp Ser Gly Thr Lys Ser Asp Met 265 270

Leu Lys Glu Leu Leu Glu IIe Asn Gln Lys Asp Glu Ser Glu Leu Ser $_{\rm et, et, tot}$

Phe Asp Glu IIe Lys His Leu Leu Leu Asp Leu Leu Val Ala Gly Thr.... 300

Asp Thr Thr Ser: Vall Thr Val Glu Trp Ala Met Thr Glu Leu Val Arg 305 310 315

Gly Leu Asn Lys Glu IIe Gln Glu Seri Asp. IIe Seri Arg Leu Pro Tyr, 340 345

Leu Arg Ala Val Val Lys Glu Ser, Phe Arg Leu His Pro Ala Thr Pro 355 360 365

Leu Ser Val Pro His Lyst Ala Asp Glu Glu Ala Glu He Asp Gly Tyr, 375 380 380 380

Ile Val Pro Lys Glý Ala Gln Val Leu Met Asn Val Trp Ala Ile Gly 1996 395 400

Arg Asp Ser Ser He Trp Arg Asn Pro Asp Val Phe Met Pro Glu Arg 405 011 415 415

Phe Leu Glu: Thr. Glu He Asp. Val Arg. Gly Gln His. Phe Glu Leu Leu 420 425 425 430

Pro Phe Gly Gly Arg Arg Lie Cys Val Gly Leu Pro Leu Ala Tyr 4445

Arg Met IIe His Leu Val Leu Ala Thr Phe IIe Ser Asp Tyr Asp Trp 450

Lys Leu Glu Gly Gly Leu Lys Thr Glu Glu Met Asp Met Ser Glu Lys 465 470 475 480

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Lys Ile

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Val Phe Asn Gln Leu Leu Asn Phe Tyr Arg Leu His Asp Tyr Met Ala 50 60

Asp Leu Ala Gly Lys Tyr Lys Thr Tyr Arg Leu Ile Ala Pro Phe Arg 65 . 70 . 75 . 80

Thr Glu Val Tyr Thr Ser Asp Pro Ala Asn Val Glu His Met Leu Lys 85 90 95

Thr Asn Phe Glu Ser Tyr Gly Lys Gly Pro Tyr Asn Cys Ser Ile Leu 100 105 110

Gly Asp Leu Phe Gly Glu Gly-Ile Phe Ala Ile Asp Gly His Lys Trp

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Arg	Glu (130×	Gln	Arg	Lys	Val	Ser 135	Ser	Leu	Glu	Phe	Ser 140	l'hr	Arg:	Val	Leu	•		MN Project	.,
Arg 145	$Asp_{()_1}^{\Sigma}$	Tyr	Ser	Ser	He 150	Val⊃ eeaf	Phe	Arg	Lys	Asn	Ala	Val	Arg	Leu	Ala 160		: 72 7)	कराह्य करता	٠
Lys	He	Leu	Ser	Gly 165:	Ala:	Alan Jana	Thr	Ser	Asn 170	Gln Ed L	Pro	Val	Asp "I i	lle 175	Gln " makeae		ingan Masara	94 () 94 () 3	i. M
Asp	Leu	Phe III	Met 180	Lỳs	Ser	Thr3 gaile	Phe	Asp 185	Sert	l·le ∷∷	Sers Hereit	Glu:	Val: 190:	Ala	nome Leu of the	1 W 1 97		eryanya Ariyanya	<u> </u>
Gly	الإغراب Val	Glu" 195	Leu	Asp	Ser	Leus	Gly: 200:	Gly	Sera Sera	Asn	Glus ∆og∉	Glu: 205	Gly	Ala	Lys	\$1. \$1)9; { } {64***	3531 4354	4
Phe	Ser. 210	lle	Ala Baa	Ala	Asp	Asp 215≥	Val 地震等	Ser:	Met	Arg 选净:	Thr 220	Leu	Trp.	Arg	Tyr.:	, ji 1 60	leigh Real	ing.	11) (6)
Val 225	Asp	Va P	Leu	Trp	Lys' 230	Leu	Lys	Arg	Alas	Leu 235	Asn	۷a l	Glya Gener	Ser	Glu 1 240	* * * * * * * * * * * * * * * * * * *	coor Cari	ur i m Fi	78 - 1
Ala	Lys	Leu	Lys	Lys 245	Ser	Leu	Gln	Val	Val 250	Asp	Glu	Phe	Val	Tyr 255	System Lys	(1034) 1. 1811	56741 17	นุกรส (,
Leu	lle	His	Ser 260	Arg	Thr	Gln	Gln	Met 265	Asn	Met	Pro	Gly	Asn 270	Asp	Ser			1	
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Lys	Gly	Asp	Met	Val 405	Ala	Tyr	Gln	Pro	Tyr 410		Met	Gly	Arg	Met 415	Lys				
Ser	lle	Trp	Gly	Asn	Asp	Ala	Glu	Glu	Phe	Lys	Pro	Glu	Arg	Trp	Leu	•			

								25/35	,						
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Gln M 465	et Lys	lle	Phe	Ser 470	Ala	lle	Leu	Leu	Arg 475	Phe	Phe	Thr	Met	Lys 480	
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Leu I	le Asp	Gly 500	Gly	Leu	Ile	Val	Arg 505		Phe	His	Arg	Met 510	Asp	Glu _.	• • • •
Lys T	hr Ala 515		:	• :	;}		• . •	•		*	. '	,			
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	34							: 1			· ;		e <u>1.</u> m z	. ' *	2 (1) 31
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<400> 63

The Europe of the Authority

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Ala Ser Asn Pro Gly Glu Ile Phe Pro Ile Leu Arg Trp Leu Gly Phe Asn Gly Leu Glu Lys Lys Leu Ala Val His Ala Arg Lys Thr Asp Glu 245 250 255 Phe Met Gln Gly Leu Leu Asp Glu His Arg Arg Gly Glu Arg Gln Asn 260 270 Thr Met Val Asp His Leu Leu Ser Leu Gln Glu Ser Gln Pro Glu Tyr 275 285 Tyr Thr Asp Glu Ile Ile Thr Gly Leu Ile Val Ala Leu Ile Ile Ala 290 295 300 Gly Thr Asp Ala Ser Val Val Thr Thr Glu Trp Ala Met Ser Leu Ile 305 310 315 Leu Asn His Pro Gln Val Leu Glu Lys Ala Arg Lys Glu Leu Asp Thr 325 335 Leu Val Gly His Glu Arg Met Val Asp Glu His Asp Leu Pro Lys Leu 340 345 350 Arg Tyr Leu His Cys Ile Val Leu Glu Thr Leu Arg Leu Phe Pro Ser. 355 Val Pro Thr Leu Val Pro His Glu Pro Ser Glu Asp Cys Lys Fle Gly 370 Gly Tyr Asn Val Pro Lys Gly Thr Met IIe Leu Val Asn Ala Trp Ala 395 Ile His Arg Asp Pro Lys Val Trp Asp Asp Pro Leu Ser Phe Lys Pro 405 410 415 Asp Arg Phe Glu Thr Met Glu Val Glu Thr His Lys Leu Leu Pro Phe Gly Met Gly Arg Arg Ala Cys Pro Gly Ala Gly Leu Ala Gln Lys Phe 435 440 445 Val Gly Leu Ala Leu Gly Ser Leu Ile Gln Cys Phe Glu Trp Glu Arg 450 455 460 Met Ser Ala Glu Lys Ile Asp Leu Asn Glu Gly Ser Gly Ile Thr Leu 465 470 475 480 Pro Lys Ala Lys Thr Leu Glu Ala Met Cys Lys Pro Arg His Ile Met 495 Glu Arg Val Leu Arg Gln Val Ser Asn Val 500

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24 00000

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